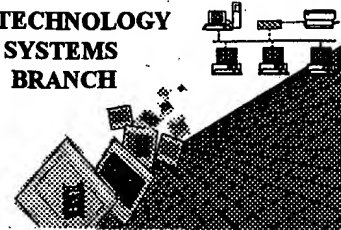


to Nelson

9B9

BIOTECHNOLOGY  
SYSTEMS  
BRANCH



TECH CENTER 1600/2900

MAY 06 2002

RECEIVED

## RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/424,091A  
Source: 1600 RUSH  
Date Processed by STIC: 4/29/2002

PT# 19

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:  
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7<sup>th</sup> Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202  
Or  
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002

# Raw Sequence Listing Error Summary

TECH CENTER  
800/2900

MAY 06 2002

RECEIVED

## ERROR DETECTED

## SUGGESTED CORRECTION

SERIAL NUMBER:

09/424,091A

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1      Wrapped Nucleics  
    Wrapped Aminos      The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2      Invalid Line Length      The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3      Misaligned Amino  
    Numbering      The numbering under each 5<sup>th</sup> amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4      Non-ASCII      The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5      Variable Length      Sequence(s)      contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6      PatentIn 2.0  
    "bug"      A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s)     . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7      Skipped Sequences  
    (OLD RULES)      Sequence(s)      missing. If intentional, please insert the following lines for each skipped sequence:  
    (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  
    (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)  
    (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  
    This sequence is intentionally skipped  
  
    Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8      Skipped Sequences  
    (NEW RULES)      Sequence(s)      missing. If intentional, please insert the following lines for each skipped sequence.  
    <210> sequence id number  
    <400> sequence id number  
    000
- 9      Use of n's or Xaa's  
    (NEW RULES)      Use of n's and/or Xaa's have been detected in the Sequence Listing.  
    Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.  
    In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 ✓ Invalid <213>  
    Response      Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 ✓ Use of <220>      Sequence(s) 1-2 missing the <220> "Feature" and associated numeric identifiers and responses.  
    Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.  
    (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12      PatentIn 2.0  
    "bug"      Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13      Misuse of n      n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.



1600

## RAW SEQUENCE LISTING

DATE: 04/29/2002

PATENT APPLICATION: US/09/424,091A

TIME: 09:49:57

Input Set : A:\#318788 v1 - 350013-66 Sequence Listing(ASCII).txt

Output Set: N:\CRF3\04292002\I424091A.raw

pp 1, 6  
Does Not Comply  
Corrected Diskette Needed

2 <110> APPLICANT: Richard Andrew Kay  
W--> 3 <120> TITLE OF INVENTION: Immunological method  
W--> 4 <130> FILE REFERENCE: DUNW/P19095US  
W--> 5 <140> CURRENT APPLICATION NUMBER: 09/424091A  
C--> 6 <141> CURRENT FILING DATE: 1999-11-09  
7 <150> PRIOR APPLICATION NUMBER: GB 9710820.3  
W--> 8 <151> PRIOR FILING DATE: 27 May 1997 1997-05-27 ← use this date format  
W--> 9 <160> NUMBER OF SEQ ID: 47

10 &lt;170&gt; SOFTWARE: SeqWin99

W--&gt; 11 &lt;210&gt; SEQ ID NO: 1

12 &lt;211&gt; LENGTH: 20

13 &lt;212&gt; TYPE: DNA

14 &lt;213&gt; ORGANISM: Artificial Sequence

W--&gt; 15 &lt;220&gt; FEATURE:

16 &lt;223&gt; OTHER INFORMATION: → see item 11 on Eur summary sheet - mandatory response needed

W--&gt; 17 &lt;400&gt; SEQUENCE: 1

18 catcagaagc agagatctcc

20

19 &lt;210&gt; SEQ ID NO: 2

20 &lt;211&gt; LENGTH: 20

21 &lt;212&gt; TYPE: DNA

22 &lt;213&gt; ORGANISM: Artificial Sequence

W--&gt; 23 &lt;220&gt; FEATURE:

24 &lt;223&gt; OTHER INFORMATION: same error

W--&gt; 25 &lt;400&gt; SEQUENCE: 2

26 gatgtcaagc tggctcgagaa

20

27 &lt;210&gt; SEQ ID NO: 3

28 &lt;211&gt; LENGTH: 18

29 &lt;212&gt; TYPE: DNA

30 &lt;213&gt; ORGANISM: Artificial Sequence

W--&gt; 31 &lt;220&gt; FEATURE:

32 &lt;223&gt; OTHER INFORMATION: 5' PCR Primer

W--&gt; 33 &lt;400&gt; SEQUENCE: 3

34 ctgaggtgca actactca

18

35 &lt;210&gt; SEQ ID NO: 4

36 &lt;211&gt; LENGTH: 24

37 &lt;212&gt; TYPE: DNA

38 &lt;213&gt; ORGANISM: Artificial Sequence

W--&gt; 39 &lt;220&gt; FEATURE:

40 &lt;223&gt; OTHER INFORMATION: 5' PCR Primer

W--&gt; 41 &lt;400&gt; SEQUENCE: 4

42 gtgttccag agggagccat tgcc

24

43 &lt;210&gt; SEQ ID NO: 5

44 &lt;211&gt; LENGTH: 21

## RAW SEQUENCE LISTING

DATE: 04/29/2002

PATENT APPLICATION: US/09/424,091A

TIME: 09:49:57

Input Set : A:\#318788 v1 - 350013-66 Sequence Listing(ASCII).txt

Output Set: N:\CRF3\04292002\I424091A.raw

45 <212> TYPE: DNA  
46 <213> ORGANISM: Artificial Sequence  
47 <220> FEATURE:  
48 <223> OTHER INFORMATION: 5' PCR Primer  
49 <400> SEQUENCE: 5  
50 ggtgaacagt caacagggag a 21  
51 <210> SEQ ID NO: 6  
52 <211> LENGTH: 21  
53 <212> TYPE: DNA  
54 <213> ORGANISM: Artificial Sequence  
55 <220> FEATURE:  
56 <223> OTHER INFORMATION: 5' PCR Primer  
57 <400> SEQUENCE: 6  
58 acaagcatta ctgtactcct a 21  
59 <210> SEQ ID NO: 7  
60 <211> LENGTH: 18  
61 <212> TYPE: DNA  
62 <213> ORGANISM: Artificial Sequence  
63 <220> FEATURE:  
64 <223> OTHER INFORMATION: 5' PCR Primer  
65 <400> SEQUENCE: 7  
66 ggccctgaac attcagga 18  
67 <210> SEQ ID NO: 8  
68 <211> LENGTH: 20  
69 <212> TYPE: DNA  
70 <213> ORGANISM: Artificial Sequence  
71 <220> FEATURE:  
72 <223> OTHER INFORMATION: 5' PCR Primer  
73 <400> SEQUENCE: 8  
74 gtcactttct agcctgctga 20  
75 <210> SEQ ID NO: 9  
76 <211> LENGTH: 21  
77 <212> TYPE: DNA  
78 <213> ORGANISM: Artificial Sequence  
79 <220> FEATURE:  
80 <223> OTHER INFORMATION: 5' PCR Primer  
81 <400> SEQUENCE: 9  
82 aggagccatt gtccagataa a 21  
83 <210> SEQ ID NO: 10  
84 <211> LENGTH: 22  
85 <212> TYPE: DNA  
86 <213> ORGANISM: Artificial Sequence  
87 <220> FEATURE:  
88 <223> OTHER INFORMATION: 5' PCR Primer  
89 <400> SEQUENCE: 10  
90 ggagagaatg tggagcagca tc 22  
91 <210> SEQ ID NO: 11  
92 <211> LENGTH: 21  
93 <212> TYPE: DNA

## RAW SEQUENCE LISTING

DATE: 04/29/2002

PATENT APPLICATION: US/09/424,091A

TIME: 09:49:57

Input Set : A:\#318788 v1 - 350013-66 Sequence Listing(ASCII).txt

Output Set: N:\CRF3\04292002\I424091A.raw

94 &lt;213&gt; ORGANISM: Artificial Sequence

95 &lt;220&gt; FEATURE:

96 &lt;223&gt; OTHER INFORMATION: 5' PCR Primer

97 &lt;400&gt; SEQUENCE: 11

98 atctcagtgc ttgtgataat a

21

99 &lt;210&gt; SEQ ID NO: 12

100 &lt;211&gt; LENGTH: 24

101 &lt;212&gt; TYPE: DNA

102 &lt;213&gt; ORGANISM: Artificial Sequence

103 &lt;220&gt; FEATURE:

104 &lt;223&gt; OTHER INFORMATION: 5' PCR Primer

105 &lt;400&gt; SEQUENCE: 12

106 acccagctgg tggagcagag ccct

24

107 &lt;210&gt; SEQ ID NO: 13

108 &lt;211&gt; LENGTH: 21

109 &lt;212&gt; TYPE: DNA

110 &lt;213&gt; ORGANISM: Artificial Sequence

111 &lt;220&gt; FEATURE:

112 &lt;223&gt; OTHER INFORMATION: 5' PCR Primer

113 &lt;400&gt; SEQUENCE: 13

114 agaaagcaag gaccaagtgt t

21

115 &lt;210&gt; SEQ ID NO: 14

116 &lt;211&gt; LENGTH: 24

117 &lt;212&gt; TYPE: DNA

118 &lt;213&gt; ORGANISM: Artificial Sequence

119 &lt;220&gt; FEATURE:

120 &lt;223&gt; OTHER INFORMATION: 5' PCR Primer

121 &lt;400&gt; SEQUENCE: 14

122 cagaaggtaa ctcaagcgca gact

24

123 &lt;210&gt; SEQ ID NO: 15

124 &lt;211&gt; LENGTH: 19

125 &lt;212&gt; TYPE: DNA

126 &lt;213&gt; ORGANISM: Artificial Sequence

127 &lt;220&gt; FEATURE:

128 &lt;223&gt; OTHER INFORMATION: 5' PCR Primer

129 &lt;400&gt; SEQUENCE: 15

130 gcttatgaga acactgcgt

19

131 &lt;210&gt; SEQ ID NO: 16

132 &lt;211&gt; LENGTH: 20

133 &lt;212&gt; TYPE: DNA

134 &lt;213&gt; ORGANISM: Artificial Sequence

135 &lt;220&gt; FEATURE:

136 &lt;223&gt; OTHER INFORMATION: 5' PCR Primer

137 &lt;400&gt; SEQUENCE: 16

138 gcagcttccc ttccagcaat

20

139 &lt;210&gt; SEQ ID NO: 17

140 &lt;211&gt; LENGTH: 20

141 &lt;212&gt; TYPE: DNA

142 &lt;213&gt; ORGANISM: Artificial Sequence

## RAW SEQUENCE LISTING

DATE: 04/29/2002

PATENT APPLICATION: US/09/424,091A

TIME: 09:49:57

Input Set : A:\#318788 v1 - 350013-66 Sequence Listing(ASCII).txt

Output Set: N:\CRF3\04292002\I424091A.raw

W--> 143 <220> FEATURE:  
 144 <223> OTHER INFORMATION: 5' PCR Primer  
 W--> 145 <400> SEQUENCE: 17  
 146 agaacctgac tgcccaggaa 20  
 147 <210> SEQ ID NO: 18  
 148 <211> LENGTH: 21  
 149 <212> TYPE: DNA  
 150 <213> ORGANISM: Artificial Sequence  
 W--> 151 <220> FEATURE:  
 152 <223> OTHER INFORMATION: 5' PCR Primer  
 W--> 153 <400> SEQUENCE: 18  
 154 catctccatg gactcatatg a 21  
 155 <210> SEQ ID NO: 19  
 156 <211> LENGTH: 19  
 157 <212> TYPE: DNA  
 158 <213> ORGANISM: Artificial Sequence  
 W--> 159 <220> FEATURE:  
 160 <223> OTHER INFORMATION: 5' PCR Primer  
 W--> 161 <400> SEQUENCE: 19  
 162 gactatacta acagcatgt 19  
 163 <210> SEQ ID NO: 20  
 164 <211> LENGTH: 18  
 165 <212> TYPE: DNA  
 166 <213> ORGANISM: Artificial Sequence  
 W--> 167 <220> FEATURE:  
 168 <223> OTHER INFORMATION: 5' PCR Primer  
 W--> 169 <400> SEQUENCE: 20  
 170 tgtcaggcaa tgacaagg 18  
 171 <210> SEQ ID NO: 21  
 172 <211> LENGTH: 26  
 173 <212> TYPE: DNA  
 174 <213> ORGANISM: Artificial Sequence  
 W--> 175 <220> FEATURE:  
 176 <223> OTHER INFORMATION: Antisense 3' PCR primer  
 W--> 177 <400> SEQUENCE: 21  
 178 aataggtcga gacacttgct actgga 26  
 179 <210> SEQ ID NO: 22  
 180 <211> LENGTH: 29  
 181 <212> TYPE: DNA  
 182 <213> ORGANISM: Artificial Sequence  
 W--> 183 <220> FEATURE:  
 184 <223> OTHER INFORMATION: Antisense mid PCR primer  
 W--> 185 <400> SEQUENCE: 22  
 186 cttgtcactg gatttagatc tctcagctg 29  
 187 <210> SEQ ID NO: 23  
 188 <211> LENGTH: 30  
 189 <212> TYPE: DNA  
 190 <213> ORGANISM: Artificial Sequence  
 W--> 191 <220> FEATURE:

## RAW SEQUENCE LISTING

DATE: 04/29/2002

PATENT APPLICATION: US/09/424,091A

TIME: 09:49:57

Input Set : A:\#318788 v1 - 350013-66 Sequence Listing(ASCII).txt

Output Set: N:\CRF3\04292002\I424091A.raw

OK 192 <223> OTHER INFORMATION: Antisense 5' PCR primer  
W-L> 193 <400> SEQUENCE: 23  
194 gtacacggca gggtcagggt tctggatatt 30  
195 <210> SEQ ID NO: 24  
196 <211> LENGTH: 30  
197 <212> TYPE: DNA  
198 <213> ORGANISM: Artificial Sequence  
OK 199 <220> FEATURE:  
200 <223> OTHER INFORMATION: 5' PCR Primer  
W--> 201 <400> SEQUENCE: 24  
202 aagagagagc aaaaggaaac attcttgaac 30  
203 <210> SEQ ID NO: 25  
204 <211> LENGTH: 30  
205 <212> TYPE: DNA  
206 <213> ORGANISM: Artificial Sequence  
W-L> 207 <220> FEATURE:  
208 <223> OTHER INFORMATION: 5' PCR Primer  
W--> 209 <400> SEQUENCE: 25  
210 gctgcaaggc cacatacgag caaggcgtcg 30  
211 <210> SEQ ID NO: 26  
212 <211> LENGTH: 30  
213 <212> TYPE: DNA  
214 <213> ORGANISM: Artificial Sequence  
OK 215 <220> FEATURE:  
216 <223> OTHER INFORMATION: 5' PCR Primer  
W-L> 217 <400> SEQUENCE: 26  
218 aaaatgaaag aaaaaggaga tattcctgag 30  
219 <210> SEQ ID NO: 27  
220 <211> LENGTH: 30  
221 <212> TYPE: DNA  
222 <213> ORGANISM: Artificial Sequence  
W-A> 223 <220> FEATURE:  
224 <223> OTHER INFORMATION: 5' PCR Primer  
W-L> 225 <400> SEQUENCE: 27  
226 ctgaggccac atatgagagt ggatttgtca 30  
227 <210> SEQ ID NO: 28  
228 <211> LENGTH: 30  
229 <212> TYPE: DNA  
230 <213> ORGANISM: Artificial Sequence  
OK 231 <220> FEATURE:  
232 <223> OTHER INFORMATION: 5' PCR Primer  
W--> 233 <400> SEQUENCE: 28  
234 cagagaaaca aaggaaactt ccctggtcga 30  
235 <210> SEQ ID NO: 29  
236 <211> LENGTH: 30  
237 <212> TYPE: DNA  
238 <213> ORGANISM: Artificial Sequence  
OK 239 <220> FEATURE:  
240 <223> OTHER INFORMATION: 5' PCR Primer

<210> 47  
<211> 30  
<212> DNA  
<213> Antisense 5' PCR primer  
<400> 47

*see item 10 on Eva Summary Sheet*

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30



## VERIFICATION SUMMARY

DATE: 04/29/2002

PATENT APPLICATION: US/09/424,091A

TIME: 09:49:58

Input Set : A:\#318788 v1 - 350013-66 Sequence Listing(ASCII).txt

Output Set: N:\CRF3\04292002\I424091A.raw

L:3 M:283 W: Missing Blank Line separator, <120> field identifier  
L:4 M:283 W: Missing Blank Line separator, <130> field identifier  
L:5 M:283 W: Missing Blank Line separator, <140> field identifier  
L:6 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:8 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD  
L:9 M:283 W: Missing Blank Line separator, <160> field identifier  
L:11 M:283 W: Missing Blank Line separator, <210> field identifier  
L:15 M:283 W: Missing Blank Line separator, <220> field identifier  
L:17 M:283 W: Missing Blank Line separator, <400> field identifier  
L:23 M:283 W: Missing Blank Line separator, <220> field identifier  
L:25 M:283 W: Missing Blank Line separator, <400> field identifier  
L:31 M:283 W: Missing Blank Line separator, <220> field identifier  
L:33 M:283 W: Missing Blank Line separator, <400> field identifier  
L:39 M:283 W: Missing Blank Line separator, <220> field identifier  
L:41 M:283 W: Missing Blank Line separator, <400> field identifier  
L:47 M:283 W: Missing Blank Line separator, <220> field identifier  
L:49 M:283 W: Missing Blank Line separator, <400> field identifier  
L:55 M:283 W: Missing Blank Line separator, <220> field identifier  
L:57 M:283 W: Missing Blank Line separator, <400> field identifier  
L:63 M:283 W: Missing Blank Line separator, <220> field identifier  
L:65 M:283 W: Missing Blank Line separator, <400> field identifier  
L:71 M:283 W: Missing Blank Line separator, <220> field identifier  
L:73 M:283 W: Missing Blank Line separator, <400> field identifier  
L:79 M:283 W: Missing Blank Line separator, <220> field identifier  
L:81 M:283 W: Missing Blank Line separator, <400> field identifier  
L:87 M:283 W: Missing Blank Line separator, <220> field identifier  
L:89 M:283 W: Missing Blank Line separator, <400> field identifier  
L:95 M:283 W: Missing Blank Line separator, <220> field identifier  
L:97 M:283 W: Missing Blank Line separator, <400> field identifier  
L:103 M:283 W: Missing Blank Line separator, <220> field identifier  
L:105 M:283 W: Missing Blank Line separator, <400> field identifier  
L:111 M:283 W: Missing Blank Line separator, <220> field identifier  
L:113 M:283 W: Missing Blank Line separator, <400> field identifier  
L:119 M:283 W: Missing Blank Line separator, <220> field identifier  
L:121 M:283 W: Missing Blank Line separator, <400> field identifier  
L:127 M:283 W: Missing Blank Line separator, <220> field identifier  
L:129 M:283 W: Missing Blank Line separator, <400> field identifier  
L:135 M:283 W: Missing Blank Line separator, <220> field identifier  
L:137 M:283 W: Missing Blank Line separator, <400> field identifier  
L:143 M:283 W: Missing Blank Line separator, <220> field identifier  
L:145 M:283 W: Missing Blank Line separator, <400> field identifier  
L:151 M:283 W: Missing Blank Line separator, <220> field identifier  
L:153 M:283 W: Missing Blank Line separator, <400> field identifier  
L:159 M:283 W: Missing Blank Line separator, <220> field identifier  
L:161 M:283 W: Missing Blank Line separator, <400> field identifier  
L:167 M:283 W: Missing Blank Line separator, <220> field identifier  
L:169 M:283 W: Missing Blank Line separator, <400> field identifier  
L:175 M:283 W: Missing Blank Line separator, <220> field identifier

**VERIFICATION SUMMARY**

PATENT APPLICATION: US/09/424,091A

DATE: 04/29/2002

TIME: 09:49:58

Input Set : A:\#318788 v1 - 350013-66 Sequence Listing(ASCII).txt

Output Set: N:\CRF3\04292002\I424091A.raw

L:177 M:283 W: Missing Blank Line separator, <400> field identifier  
L:183 M:283 W: Missing Blank Line separator, <220> field identifier  
L:185 M:283 W: Missing Blank Line separator, <400> field identifier  
L:191 M:283 W: Missing Blank Line separator, <220> field identifier